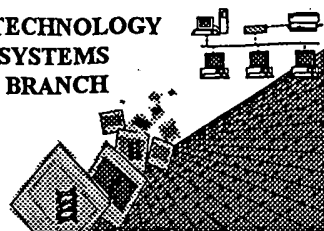


BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/099,700
Source: OIPE
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



Does Not Comply
Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/099,700

DATE: 04/03/2002
TIME: 14:38:32

Input Set : A:\1613seq.001
Output Set: N:\CRF3\04032002\J099700.raw

3 <110> APPLICANT: Edwin L. Madison
4 Edgar O. Ong
6 <120> TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING TRANSMEMBRANE SERINE
PROTEASE 7, THE
7 ENCODED POLYPEPTIDES AND METHODS BASED THEREON
9 <130> FILE REFERENCE: 24745-1613
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/099,700
C--> 12 <141> CURRENT FILING DATE: 2002-03-13
14 <150> PRIOR APPLICATION NUMBER: 60/275,592
15 <151> PRIOR FILING DATE: 2001-03-13
17 <160> NUMBER OF SEQ ID NOS: 22
19 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

983 <210> SEQ ID NO: 21
984 <211> LENGTH: (35) → Input 35, counted 36!
985 <212> TYPE: DNA
986 <213> ORGANISM: Artificial Sequence
988 <220> FEATURE:
989 <223> OTHER INFORMATION: Primer
991 <400> SEQUENCE: 21
E--> 992 attcgcgcc gcctacatac cagtctttga ggcaat

(35)

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/099,700

DATE: 04/03/2002

TIME: 14:38:33

Input Set : A:\1613seq.001

Output Set: N:\CRF3\04032002\J099700.raw

L:11 M:270 C: Current Application Number differs, Replaced Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:992 M:254 E: No. of Bases conflict, LENGTH:Input:35 Counted:36 SEQ:21

L:992 M:252 E: No. of Seq. differs, <211>LENGTH:Input:35 Found:36 SEQ:21